

Patent Application US/08/062,021

1 SEQUENCE LISTING
2

3 (1) GENERAL INFORMATION
4 (i) APPLICANT: Lynn Bergmeyer
5 Thomas J. Cummins
6 John B. Findlay
7 JoAnne H. Kerschner
8 (ii) TITLE OF THE INVENTION: DIAGNOSTIC
9 COMPOSITIONS, ELEMENTS, METHODS AND TEST KITS
10 FOR AMPLIFICATION AND DETECTION OF HUMAN CMV
11 DNA USING PRIMERS HAVING MATCHED MELTING
12 TEMPERATURES
13 (iii) NUMBER OF SEQUENCES: 26
14 (iv) CORRESPONDENCE ADDRESS:
15 (A) ADDRESSEE: Eastman Kodak Company,
16 Patent Legal Staff
17 (B) STREET: 343 State Street
18 (C) CITY: Rochester
19 (D) STATE: New York
20 (E) COUNTRY: U.S.A.
21 (F) ZIP: 14650 - 2201
22 (v) COMPUTER READABLE FORM:
23 (A) MEDIUM TYPE: Diskette, 3.5
24 inch, 1.44 MB storage (IBM)
25 (B) COMPUTER: IBM PS/2
26 (C) OPERATING SYSTEM: MS-DOS Version
27 3.3
28 (D) SOFTWARE: PC-8 (Word for Windows)
29 (vi) CURRENT APPLICATION DATA:
30 (A) APPLICATION NUMBER: To Be Assigned
31 (B) FILING DATE: To Be Assigned
32 (C) CLASSIFICATION: To Be Assigned
33 (vii) PRIOR APPLICATION DATE: None
34 (viii) ATTORNEY/AGENT INFORMATION:
35 (A) NAME: Tucker, J. Lanny
36 (B) REGISTRATION NUMBER: 27,678
37 (C) REFERENCE/DOCKET NUMBER: 67270
38 (ix) TELECOMMUNICATION INFORMATION:
39 (A) TELEPHONE: (716) 722-9332
40 (B) TELEFAX: (716) 477-4646
41
42 (2) INFORMATION FOR SEQ ID: NO: 1
43 (i) SEQUENCE CHARACTERISTICS:
44 (A) LENGTH: 25 nucleotides
45 (B) TYPE: Nucleic acid
46 (C) STRANDEDNESS: Single
47 (D) TOPOLOGY: Linear
48 (ii) MOLECULE TYPE: Primer for hCMV DNA
49 (iii) HYPOTHETICAL: No
50 (iv) ANTI-SENSE: No
51 (vi) ORIGINAL SOURCE: Synthetically prepared
52 (vii) IMMEDIATE SOURCE: Same

See pp. 1 + 6

This colon is not correct throughout the sequence listing. Please delete this throughout the sequence listing. I suggest a "Search and replace" function for this: Search for all "ID:" & replace with "ID" + space).

Patent Application US/08/062,021

53 (x) PUBLICATION INFORMATION: U.S. 5,147,777
54 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:1
55
56 GAGGCTATTG TAGCCTACAC TTTGG 25
57
58 (3) INFORMATION FOR SEQ ID:NO:2
59 (i) SEQUENCE CHARACTERISTICS:
60 (A) LENGTH: 25 nucleotides
61 (B) TYPE: Nucleic acid
62 (C) STRANDEDNESS: Single
63 (D) TOPOLOGY: Linear
64 (ii) MOLECULE TYPE: Primer for hCMV DNA
65 (iii) HYPOTHETICAL: No
66 (iv) ANTI-SENSE: No
67 (vi) ORIGINAL SOURCE: Synthetically prepared
68 (vii) IMMEDIATE SOURCE: Same
69 (x) PUBLICATION INFORMATION: U.S. 5,147,777
70 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:2
71
72 CAGCACCATC CTCCTCTTCC TCTGG 25
73
74
75
76 (4) INFORMATION FOR SEQ ID:NO:3
77 (i) SEQUENCE CHARACTERISTICS:
78 (A) LENGTH: 25 nucleotides
79 (B) TYPE: Nucleic acid
80 (C) STRANDEDNESS: Single
81 (D) TOPOLOGY: Linear
82 (ii) MOLECULE TYPE: Primer for hCMV DNA
83 (iii) HYPOTHETICAL: No
84 (iv) ANTI-SENSE: No
85 (vi) ORIGINAL SOURCE: Synthetically prepared
86 (vii) IMMEDIATE SOURCE: Same
87 (x) PUBLICATION INFORMATION: None
88 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:3
89
90 TGCACTGCCA GGTGCTTCGG CTCAT 25
91
92 (5) INFORMATION FOR SEQ ID:NO:4
93 (i) SEQUENCE CHARACTERISTICS:
94 (A) LENGTH: 25 nucleotides
95 (B) TYPE: Nucleic acid
96 (C) STRANDEDNESS: Single
97 (D) TOPOLOGY: Linear
98 (ii) MOLECULE TYPE: Primer for hCMV DNA
99 (iii) HYPOTHETICAL: No
100 (iv) ANTI-SENSE: No
101 (vi) ORIGINAL SOURCE: Synthetically prepared
102 (vii) IMMEDIATE SOURCE: Same
103 (x) PUBLICATION INFORMATION: U.S. 5,147,777
104 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:4

Patent Application US/08/062,021

105
106 CACCACGCAG CGGCCCTTGA TGTTC 25
107
108
109
110 (6) INFORMATION FOR SEQ ID:NO:5
111 (i) SEQUENCE CHARACTERISTICS:
112 (A) LENGTH: 30 nucleotides
113 (B) TYPE: Nucleic acid
114 (C) STRANDEDNESS: Single
115 (D) TOPOLOGY: Linear
116 (ii) MOLECULE TYPE: Probe for hCMV DNA
117 (iii) HYPOTHETICAL: No
118 (iv) ANTI-SENSE: No
119 (vi) ORIGINAL SOURCE: Synthetically prepared
120 (vii) IMMEDIATE SOURCE: Same
121 (x) PUBLICATION INFORMATION: U.S. 5,147,777
122 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:5
123
124 GGTGTCACCC CCAGAGTCCC CTGTACCCGC 30
125
126 (7) INFORMATION FOR SEQ ID:NO:6
127 (i) SEQUENCE CHARACTERISTICS:
128 (A) LENGTH: 30 nucleotides
129 (B) TYPE: Nucleic acid
130 (C) STRANDEDNESS: Single
131 (D) TOPOLOGY: Linear
132 (ii) MOLECULE TYPE: Probe for hCMV DNA
133 (iii) HYPOTHETICAL: No
134 (iv) ANTI-SENSE: No
135 (vi) ORIGINAL SOURCE: Synthetically prepared
136 (vii) IMMEDIATE SOURCE: Same
137 (x) PUBLICATION INFORMATION: U.S. 5,147,777
138 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:6
139
140 GACACAGTGT CCTCCCGCTC CTCTTGAGCA 30
141
142
143
144 (8) INFORMATION FOR SEQ ID:NO:7
145 (i) SEQUENCE CHARACTERISTICS:
146 (A) LENGTH: 30 nucleotides
147 (B) TYPE: Nucleic acid
148 (C) STRANDEDNESS: Single
149 (D) TOPOLOGY: Linear
150 (ii) MOLECULE TYPE: Probe for hCMV DNA
151 (iii) HYPOTHETICAL: No
152 (iv) ANTI-SENSE: No
153 (vi) ORIGINAL SOURCE: Synthetically prepared
154 (vii) IMMEDIATE SOURCE: Same
155 (x) PUBLICATION INFORMATION: U.S. 5,147,777
156 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:7

Patent Application US/08/062,021

157
158 GTGGAAGGCG GCTCGCTGGA AGCCGGTCGT 30
159
160 (9) INFORMATION FOR SEQ ID:NO:8
161 (i) SEQUENCE CHARACTERISTICS:
162 (A) LENGTH: 30 nucleotides
163 (B) TYPE: Nucleic acid
164 (C) STRANDEDNESS: Single
165 (D) TOPOLOGY: Linear
166 (ii) MOLECULE TYPE: Probe for hCMV DNA
167 (iii) HYPOTHETICAL: No
168 (iv) ANTI-SENSE: No
169 (vi) ORIGINAL SOURCE: Synthetically prepared
170 (vii) IMMEDIATE SOURCE: Same
171 (x) PUBLICATION INFORMATION: U.S. 5,147,777
172 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:8
173
174 GAACCGAGGG CCGGCTCACC TCTATGTTGG 30
175
176
177
178 (10) INFORMATION FOR SEQ ID:NO:9
179 (i) SEQUENCE CHARACTERISTICS:
180 (A) LENGTH: 41 nucleotides
181 (B) TYPE: Nucleic acid
182 (C) STRANDEDNESS: Single
183 (D) TOPOLOGY: Linear
184 (ii) MOLECULE TYPE: Probe for HIV-I DNA
185 (iii) HYPOTHETICAL: No
186 (iv) ANTI-SENSE: No
187 (vi) ORIGINAL SOURCE: Synthetically prepared
188 (vii) IMMEDIATE SOURCE: Same
189 (x) PUBLICATION INFORMATION: Unknown
190 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:9
191
192 ATCCTGGGAT TAAATAAAAT AGTAAGAACATG TATAGCCCTA C 41
193
194 (11) INFORMATION FOR SEQ ID:NO:10
195 (i) SEQUENCE CHARACTERISTICS:
196 (A) LENGTH: 28 nucleotides
197 (B) TYPE: Nucleic acid
198 (C) STRANDEDNESS: Single
199 (D) TOPOLOGY: Linear
200 (ii) MOLECULE TYPE: Primer for HIV-I DNA
201 (iii) HYPOTHETICAL: No
202 (iv) ANTI-SENSE: No
203 (vi) ORIGINAL SOURCE: Synthetically prepared
204 (vii) IMMEDIATE SOURCE: Same
205 (x) PUBLICATION INFORMATION: Unknown
206 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:10
207
208 AGTGGGGGGA CATCAAGCAG CCATGCAA 28

Patent Application US/08/062,021

209
210
211
212 (12) INFORMATION FOR SEQ ID:NO:11
213 (i) SEQUENCE CHARACTERISTICS:
214 (A) LENGTH: 26 nucleotides
215 (B) TYPE: Nucleic acid
216 (C) STRANDEDNESS: Single
217 (D) TOPOLOGY: Linear
218 (ii) MOLECULE TYPE: Primer for HIV-I DNA
219 (iii) HYPOTHETICAL: No
220 (iv) ANTI-SENSE: No
221 (vi) ORIGINAL SOURCE: Synthetically prepared
222 (vii) IMMEDIATE SOURCE: Same
223 (x) PUBLICATION INFORMATION: None
224 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:11
225
226 CTTGGTTCTC TCATCTGGCC TGGTGC 26
227
228 (13) INFORMATION FOR SEQ ID:NO:12
229 (i) SEQUENCE CHARACTERISTICS:
230 (A) LENGTH: 28 nucleotides
231 (B) TYPE: Nucleic acid
232 (C) STRANDEDNESS: Single
233 (D) TOPOLOGY: Linear
234 (ii) MOLECULE TYPE: Primer for HIV-I DNA
235 (iii) HYPOTHETICAL: No
236 (iv) ANTI-SENSE: No
237 (vi) ORIGINAL SOURCE: Synthetically prepared
238 (vii) IMMEDIATE SOURCE: Same
239 (x) PUBLICATION INFORMATION: None
240 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:12
241
242 TAGCACCCAC CAGGGCAAAG AGAAGAGT 28
243
244
245
246 (14) INFORMATION FOR SEQ ID:NO:13
247 (i) SEQUENCE CHARACTERISTICS:
248 (A) LENGTH: 28 nucleotides
249 (B) TYPE: Nucleic acid
250 (C) STRANDEDNESS: Single
251 (D) TOPOLOGY: Linear
252 (ii) MOLECULE TYPE: Primer for HIV-I DNA
253 (iii) HYPOTHETICAL: No
254 (iv) ANTI-SENSE: No
255 (vi) ORIGINAL SOURCE: Synthetically prepared
256 (vii) IMMEDIATE SOURCE: Same
257 (x) PUBLICATION INFORMATION: None
258 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:13
259
260 AGATGCTGTT GCGCCTCAAT AGCCCTCA 28

Patent Application US/08/062,021

261
262 (15) INFORMATION FOR SEQ ID:NO:14
263 (i) SEQUENCE CHARACTERISTICS:
264 (A) LENGTH: 28 nucleotides
265 (B) TYPE: Nucleic acid
266 (C) STRANDEDNESS: Single
267 (D) TOPOLOGY: Linear
268 (ii) MOLECULE TYPE: Probe for HIV-I DNA
269 (iii) HYPOTHETICAL: No
270 (iv) ANTI-SENSE: No
271 (vi) ORIGINAL SOURCE: Synthetically prepared
272 (vii) IMMEDIATE SOURCE: Same
273 (x) PUBLICATION INFORMATION: None
274 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:14
275
276 GAGACCATCA ATGAGGAAGC TGCAGAAT 28
277
278
279
280 (16) INFORMATION FOR SEQ ID:NO:15
281 (i) SEQUENCE CHARACTERISTICS:
282 (A) LENGTH: 28 nucleotides
283 (B) TYPE: Nucleic acid
284 (C) STRANDEDNESS: Single
285 (D) TOPOLOGY: Linear
286 (ii) MOLECULE TYPE: Probe for HIV-I DNA
287 (iii) HYPOTHETICAL: No
288 (iv) ANTI-SENSE: No
289 (vi) ORIGINAL SOURCE: Synthetically prepared
290 (vii) IMMEDIATE SOURCE: Same
291 (x) PUBLICATION INFORMATION: None
292 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:15
293
294 GTGCAGCAGC AGAACAAATT GCTGAGGG 28
295
296 (17) INFORMATION FOR SEQ ID:NO:16
297 (i) SEQUENCE CHARACTERISTICS:
298 (A) LENGTH: 27 nucleotides
299 (B) TYPE: Nucleic acid
300 (C) STRANDEDNESS: Single
301 (D) TOPOLOGY: Linear
302 (ii) MOLECULE TYPE: Primer for hCMV DNA
303 (iii) HYPOTHETICAL: No
304 (iv) ANTI-SENSE: No
305 (vi) ORIGINAL SOURCE: Synthetically prepared
306 (vii) IMMEDIATE SOURCE: Same
307 (x) PUBLICATION INFORMATION: None
308 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:16
309
310 5' CATTCCCACT GACTTTCTGA CGCACGT-3' 27
311
312

delete -
(Sequences 1-15
are correct.
Sequences 16-26
are not because
of this).

Patent Application US/08/062,021

313
314 (18) INFORMATION FOR SEQ ID:NO:17
315 (i) SEQUENCE CHARACTERISTICS:
316 (A) LENGTH: 24 nucleotides
317 (B) TYPE: Nucleic acid
318 (C) STRANDEDNESS: Single
319 (D) TOPOLOGY: Linear
320 (ii) MOLECULE TYPE: Primer for hCMV DNA
321 (iii) HYPOTHETICAL: No
322 (iv) ANTI-SENSE: No
323 (vi) ORIGINAL SOURCE: Synthetically prepared
324 (vii) IMMEDIATE SOURCE: Same
325 (x) PUBLICATION INFORMATION: None
326 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:17
327
328 5'-TGAGGTCGTG GAACTTGATG GCGT-3' 24
329
330 (19) INFORMATION FOR SEQ ID:NO:18
331 (i) SEQUENCE CHARACTERISTICS:
332 (A) LENGTH: 30 nucleotides
333 (B) TYPE: Nucleic acid
334 (C) STRANDEDNESS: Single
335 (D) TOPOLOGY: Linear
336 (ii) MOLECULE TYPE: Probe for hCMV DNA
337 (iii) HYPOTHETICAL: No
338 (iv) ANTI-SENSE: No
339 (vi) ORIGINAL SOURCE: Synthetically prepared
340 (vii) IMMEDIATE SOURCE: Same
341 (x) PUBLICATION INFORMATION: None
342 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:18
343
344 5'-GGTCATCGCC GTAGTAGATG CGTAAGGCCT-3' 30
345
346
347
348 (20) INFORMATION FOR SEQ ID:NO:19
349 (i) SEQUENCE CHARACTERISTICS:
350 (A) LENGTH: 26 nucleotides
351 (B) TYPE: Nucleic acid
352 (C) STRANDEDNESS: Single
353 (D) TOPOLOGY: Linear
354 (ii) MOLECULE TYPE: Primer for avian
355 endogenous
356 provirus ev-1 DNA
357 (iii) HYPOTHETICAL: No
358 (iv) ANTI-SENSE: No
359 (vi) ORIGINAL SOURCE: Synthetically prepared
360 (vii) IMMEDIATE SOURCE: Same
361 (x) PUBLICATION INFORMATION: None
362 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:19
363
364 5'-GGAATGACGC AAGGACATAT GGGCGT-3' 26

Patent Application US/08/062,021

365
366 (21) INFORMATION FOR SEQ ID:NO:20
367 (i) SEQUENCE CHARACTERISTICS:
368 (A) LENGTH: 26 nucleotides
369 (B) TYPE: Nucleic acid
370 (C) STRANDEDNESS: Single
371 (D) TOPOLOGY: Linear
372 (ii) MOLECULE TYPE: Primer for avian
373 endogenous
374 provirus ev-1 DNA
375 (iii) HYPOTHETICAL: No
376 (iv) ANTI-SENSE: No
377 (vi) ORIGINAL SOURCE: Synthetically prepared
378 (vii) IMMEDIATE SOURCE: Same
379 (x) PUBLICATION INFORMATION: None
380 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:20
381
382 5'-CCCAAGGTGCA CACCAATGTG GTGGAT-3' 26
383
384
385
386 (22) INFORMATION FOR SEQ ID:NO:21
387 (i) SEQUENCE CHARACTERISTICS:
388 (A) LENGTH: 25 nucleotides
389 (B) TYPE: Nucleic acid
390 (C) STRANDEDNESS: Single
391 (D) TOPOLOGY: Linear
392 (ii) MOLECULE TYPE: Primer for positive
control target DNA
394 (iii) HYPOTHETICAL: No
395 (iv) ANTI-SENSE: No
396 (vi) ORIGINAL SOURCE: Synthetically prepared
397 (vii) IMMEDIATE SOURCE: Same
398 (x) PUBLICATION INFORMATION: None
399 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:21
400
401 5'-GGACTGTGCG CGTTGTATAAC CCTGC-3' 25
402
403 (23) INFORMATION FOR SEQ ID:NO:22
404 (i) SEQUENCE CHARACTERISTICS:
405 (A) LENGTH: 25 nucleotides
406 (B) TYPE: Nucleic acid
407 (C) STRANDEDNESS: Single
408 (D) TOPOLOGY: Linear
409 (ii) MOLECULE TYPE: Primer for positive
control target DNA
411 (iii) HYPOTHETICAL: No
412 (iv) ANTI-SENSE: No
413 (vi) ORIGINAL SOURCE: Synthetically prepared
414 (vii) IMMEDIATE SOURCE: Same
415 (x) PUBLICATION INFORMATION: None
416 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:22

Patent Application US/08/062,021

417
418 5'-ACTCCCGAAG CGAATGGCAC GTGGA-3' 25
419
420
421

422 (24) INFORMATION FOR SEQ ID:NO:23
423 (i) SEQUENCE CHARACTERISTICS:
424 (A) LENGTH: 25 nucleotides
425 (B) TYPE: Nucleic acid
426 (C) STRANDEDNESS: Single
427 (D) TOPOLOGY: Linear
428 (ii) MOLECULE TYPE: Primer for positive
control target DNA
430 (iii) HYPOTHETICAL: No
431 (iv) ANTI-SENSE: No
432 (vi) ORIGINAL SOURCE: Synthetically prepared
433 (vii) IMMEDIATE SOURCE: Same
434 (x) PUBLICATION INFORMATION: None
435 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:23
436

437 5'-CATAGCTTGT GCCCGTGTGG CACGT-3' 25
438

439 (25) INFORMATION FOR SEQ ID:NO:24
440 (i) SEQUENCE CHARACTERISTICS:
441 (A) LENGTH: 25 nucleotides
442 (B) TYPE: Nucleic acid
443 (C) STRANDEDNESS: Single
444 (D) TOPOLOGY: Linear
445 (ii) MOLECULE TYPE: Primer for positive
control target DNA
447 (iii) HYPOTHETICAL: No
448 (iv) ANTI-SENSE: No
449 (vi) ORIGINAL SOURCE: Synthetically prepared
450 (vii) IMMEDIATE SOURCE: Same
451 (x) PUBLICATION INFORMATION: None
452 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:24
453

454 5'-CCAAGACGAG ACCGTCAGAG CTGGT-3' 25
455
456
457

458 (26) INFORMATION FOR SEQ ID:NO:25
459 (i) SEQUENCE CHARACTERISTICS:
460 (A) LENGTH: 26 nucleotides
461 (B) TYPE: Nucleic acid
462 (C) STRANDEDNESS: Single
463 (D) TOPOLOGY: Linear
464 (ii) MOLECULE TYPE: Probe for avian
endogenous
466 provirus ev-1 DNA
467 (iii) HYPOTHETICAL: No
468 (iv) ANTI-SENSE: No

Patent Application US/08/062,021

469 (vi) ORIGINAL SOURCE: Synthetically prepared
470 (vii) IMMEDIATE SOURCE: Same
471 (x) PUBLICATION INFORMATION: None
472 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:25
473
474 5'-AAGCTGTTGC CGCCATCAA TAAACG-3' 26
475
476 (27) INFORMATION FOR SEQ ID:NO:26
477 (i) SEQUENCE CHARACTERISTICS:
478 (A) LENGTH: 30 nucleotides
479 (B) TYPE: Nucleic acid
480 (C) STRANDEDNESS: Single
481 (D) TOPOLOGY: Linear
482 (ii) MOLECULE TYPE: Probe for positive
control target DNA
484 (iii) HYPOTHETICAL: No
485 (iv) ANTI-SENSE: No
486 (vi) ORIGINAL SOURCE: Synthetically prepared
487 (vii) IMMEDIATE SOURCE: Same
488 (x) PUBLICATION INFORMATION: None
489 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:26
490
491 5'-CTGCGTTAGA CCGAGAACTG TGGATAAAGG-3' 30
492

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/062,021

DATE: 06/04/93
TIME: 15:24:00
S5726

LINE ERROR

ORIGINAL TEXT

24 Response Exceeds Line Limitations
27 Response Exceeds Line Limitations
30 Wrong application Serial Number
32 Wrong Classification
54 Wrong Sequence Number
70 Wrong Sequence Number
88 Wrong Sequence Number
104 Wrong Sequence Number
122 Wrong Sequence Number
138 Wrong Sequence Number
156 Wrong Sequence Number
172 Wrong Sequence Number
190 Wrong Sequence Number
206 Wrong Sequence Number
224 Wrong Sequence Number
240 Wrong Sequence Number
258 Wrong Sequence Number
274 Wrong Sequence Number
292 Wrong Sequence Number
308 Wrong Sequence Number
310 Wrong Nucleic Acid Designator (5)
310 Wrong Nucleic Acid Designator (')
310 Wrong Nucleic Acid Designator (-)
310 Wrong Nucleic Acid Designator (-)
310 Wrong Nucleic Acid Designator (3)
310 Wrong Nucleic Acid Designator (')
326 Wrong Sequence Number
328 Wrong Nucleic Acid Designator (5)
328 Wrong Nucleic Acid Designator (')
328 Wrong Nucleic Acid Designator (-)
328 Wrong Nucleic Acid Designator (-)
328 Wrong Nucleic Acid Designator (3)
328 Wrong Nucleic Acid Designator (')
342 Wrong Sequence Number
344 Wrong Nucleic Acid Designator (5)
344 Wrong Nucleic Acid Designator (')
344 Wrong Nucleic Acid Designator (-)
344 Wrong Nucleic Acid Designator (-)
344 Wrong Nucleic Acid Designator (3)
344 Wrong Nucleic Acid Designator (')
362 Wrong Sequence Number
364 Wrong Nucleic Acid Designator (5)
364 Wrong Nucleic Acid Designator (')
364 Wrong Nucleic Acid Designator (-)
364 Wrong Nucleic Acid Designator (-)
364 Wrong Nucleic Acid Designator (3)
364 Wrong Nucleic Acid Designator (')
380 Wrong Sequence Number
382 Wrong Nucleic Acid Designator (5)
382 Wrong Nucleic Acid Designator (')
382 Wrong Nucleic Acid Designator (-)
382 Wrong Nucleic Acid Designator (-)

inch, 1.44 MB storage (IBM)
3.3
(A) APPLICATION NUMBER: To Be Assigned
(C) CLASSIFICATION: To Be Assigned
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 1
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 2
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 3
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 4
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 5
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 6
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 7
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 8
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 9
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 10
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 11
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 12
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 13
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 14
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 15
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 16
5'-CATTCCCACT GACTTCTGA CGCACGT-3' 27
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 17
5'-TGAGGTCGTG GAACTTGATG GCGT-3' 24
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 18
5'-GGTCATGCC GTAGTAGATG CGTAAGGCCT-3'
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 19
5'-GGAATGACGC AAGGACATAT GGGCGT-3' 26
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO: 20
5'-CCCAGGTGCA CACCAATGTG GTGGAT-3' 26
5'-CCCAGGTGCA CACCAATGTG GTGGAT-3' 26
5'-CCCAGGTGCA CACCAATGTG GTGGAT-3' 26
5'-CCCAGGTGCA CACCAATGTG GTGGAT-3' 26

LINE ERROR

382	Wrong	Nucleic Acid Designator	(3)
382	Wrong	Nucleic Acid Designator	(')
399	Wrong	Sequence Number	
401	Wrong	Nucleic Acid Designator	(5)
401	Wrong	Nucleic Acid Designator	(')
401	Wrong	Nucleic Acid Designator	(-)
401	Wrong	Nucleic Acid Designator	(-)
401	Wrong	Nucleic Acid Designator	(3)
401	Wrong	Nucleic Acid Designator	(')
416	Wrong	Sequence Number	
418	Wrong	Nucleic Acid Designator	(5)
418	Wrong	Nucleic Acid Designator	(')
418	Wrong	Nucleic Acid Designator	(-)
418	Wrong	Nucleic Acid Designator	(-)
418	Wrong	Nucleic Acid Designator	(3)
418	Wrong	Nucleic Acid Designator	(')
435	Wrong	Sequence Number	
437	Wrong	Nucleic Acid Designator	(5)
437	Wrong	Nucleic Acid Designator	(')
437	Wrong	Nucleic Acid Designator	(-)
437	Wrong	Nucleic Acid Designator	(-)
437	Wrong	Nucleic Acid Designator	(3)
437	Wrong	Nucleic Acid Designator	(')
452	Wrong	Sequence Number	
454	Wrong	Nucleic Acid Designator	(5)
454	Wrong	Nucleic Acid Designator	(')
454	Wrong	Nucleic Acid Designator	(-)
454	Wrong	Nucleic Acid Designator	(-)
454	Wrong	Nucleic Acid Designator	(3)
454	Wrong	Nucleic Acid Designator	(')
472	Wrong	Sequence Number	
474	Wrong	Nucleic Acid Designator	(5)
474	Wrong	Nucleic Acid Designator	(')
474	Wrong	Nucleic Acid Designator	(-)
474	Wrong	Nucleic Acid Designator	(-)
474	Wrong	Nucleic Acid Designator	(3)
474	Wrong	Nucleic Acid Designator	(')
489	Wrong	Sequence Number	
491	Wrong	Nucleic Acid Designator	(5)
491	Wrong	Nucleic Acid Designator	(')
491	Wrong	Nucleic Acid Designator	(-)
491	Wrong	Nucleic Acid Designator	(-)
491	Wrong	Nucleic Acid Designator	(3)
491	Wrong	Nucleic Acid Designator	(')

PAGE: 1

SEQUENCE MISSING ITEM REPORT
PATENT APPLICATION US/08/062,021

DATE: 06/04/93
TIME: 15:24:00
S5726

MANDATORY IDENTIFIER THAT WAS NOT FOUND

APPLICATION NUMBER
FILING DATE

PAGE: 1

SEQUENCE CORRECTION REPORT
PATENT APPLICATION US/08/062,021

DATE: 06/04/93
TIME: 15:24:00
S5726

LINE ORIGINAL TEXT

3 (1) GENERAL INFORMATION
8 (ii) TITLE OF THE INVENTION: DIAGNOSTIC
33 (vii) PRIOR APPLICATION DATE: None
42 (2) INFORMATION FOR SEQ ID: NO:1
54 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:1
58 (3) INFORMATION FOR SEQ ID: NO:2
70 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:2
76 (4) INFORMATION FOR SEQ ID: NO:3
88 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:3
92 (5) INFORMATION FOR SEQ ID: NO:4
104 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:4
110 (6) INFORMATION FOR SEQ ID: NO:5
122 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:5
126 (7) INFORMATION FOR SEQ ID: NO:6
138 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:6
144 (8) INFORMATION FOR SEQ ID: NO:7
156 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:7
160 (9) INFORMATION FOR SEQ ID: NO:8
172 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:8
178 (10) INFORMATION FOR SEQ ID: NO:9
190 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:9
194 (11) INFORMATION FOR SEQ ID: NO:10
206 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:10
212 (12) INFORMATION FOR SEQ ID: NO:11
224 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:11
228 (13) INFORMATION FOR SEQ ID: NO:12
240 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:12
246 (14) INFORMATION FOR SEQ ID: NO:13
258 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:13
262 (15) INFORMATION FOR SEQ ID: NO:14
274 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:14
280 (16) INFORMATION FOR SEQ ID: NO:15
292 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:15
296 (17) INFORMATION FOR SEQ ID: NO:16
308 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:16
314 (18) INFORMATION FOR SEQ ID: NO:17
326 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:17
330 (19) INFORMATION FOR SEQ ID: NO:18
342 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:18
348 (20) INFORMATION FOR SEQ ID: NO:19
362 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:19
366 (21) INFORMATION FOR SEQ ID: NO:20
380 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:20
386 (22) INFORMATION FOR SEQ ID: NO:21
399 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:21
403 (23) INFORMATION FOR SEQ ID: NO:22
416 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:22
422 (24) INFORMATION FOR SEQ ID: NO:23
435 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:23
439 (25) INFORMATION FOR SEQ ID: NO:24
452 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:24
458 (26) INFORMATION FOR SEQ ID: NO:25
472 (xi) SEQUENCE DESCRIPTION: SEQ ID: NO:25

CORRECTED TEXT

(1) GENERAL INFORMATION:
(ii) TITLE OF INVENTION: DIAGNOSTIC
(vii) PRIOR APPLICATION DATA: None
(2) INFORMATION FOR SEQ ID NO: NO:1:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:1:
(2) INFORMATION FOR SEQ ID NO: NO:2:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:2:
(2) INFORMATION FOR SEQ ID NO: NO:3:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:3:
(2) INFORMATION FOR SEQ ID NO: NO:4:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:4:
(2) INFORMATION FOR SEQ ID NO: NO:5:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:5:
(2) INFORMATION FOR SEQ ID NO: NO:6:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:6:
(2) INFORMATION FOR SEQ ID NO: NO:7:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:7:
(2) INFORMATION FOR SEQ ID NO: NO:8:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:8:
(2) INFORMATION FOR SEQ ID NO: NO:9:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:9:
(2) INFORMATION FOR SEQ ID NO: NO:10:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:10:
(2) INFORMATION FOR SEQ ID NO: NO:11:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:11:
(2) INFORMATION FOR SEQ ID NO: NO:12:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:12:
(2) INFORMATION FOR SEQ ID NO: NO:13:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:13:
(2) INFORMATION FOR SEQ ID NO: NO:14:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:14:
(2) INFORMATION FOR SEQ ID NO: NO:15:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:15:
(2) INFORMATION FOR SEQ ID NO: NO:16:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:16:
(2) INFORMATION FOR SEQ ID NO: NO:17:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:17:
(2) INFORMATION FOR SEQ ID NO: NO:18:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:18:
(2) INFORMATION FOR SEQ ID NO: NO:19:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:19:
(2) INFORMATION FOR SEQ ID NO: NO:20:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:20:
(2) INFORMATION FOR SEQ ID NO: NO:21:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:21:
(2) INFORMATION FOR SEQ ID NO: NO:22:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:22:
(2) INFORMATION FOR SEQ ID NO: NO:23:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:23:
(2) INFORMATION FOR SEQ ID NO: NO:24:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:24:
(2) INFORMATION FOR SEQ ID NO: NO:25:
(xi) SEQUENCE DESCRIPTION: SEQ ID: NO:25

PAGE: 2

SEQUENCE CORRECTION REPORT
PATENT APPLICATION US/08/062,021

DATE: 06/04/93
TIME: 15:24:00
S5726

LINE ORIGINAL TEXT

476 (27) INFORMATION FOR SEQ ID:NO:26
489 (xi) SEQUENCE DESCRIPTION: SEQ ID:NO:26

CORRECTED TEXT

(2) INFORMATION FOR SEQ ID NO:NO:26:
(xi) SEQUENCE DESCRIPTION: SEQ ID:NO:26